

THAT WHICH IS CLAIMED IS:

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- INNOCENCE*
- 1 38. A stably transformed duckweed plant comprising a heterologous nucleic acid of interest incorporated in its genome.
- 2 40. The stably transformed duckweed plant according to Claim 38, wherein said duckweed plant comprises fewer than 5 copies of said heterologous nucleic acid of interest.
- 3 41. The stably transformed duckweed plant according to Claim 38, wherein said duckweed plant is selected from the group consisting of the genus *Spirodela*, genus *Wolffia*, genus *Wolfiella*, and genus *Lemna*.
- 4 42. The stably transformed duckweed plant according to Claim 38, wherein said duckweed plant is selected from the genus *Lemna*.
- 15 5 43. The stably transformed duckweed plant according to Claim 38, wherein said duckweed plant is selected from the group consisting of a species of *Lemna minor*, a species of *Lemna miniscula*, and a species of *Lemna gibba*.
- 6 44. The stably transformed duckweed plant according to Claim 38, wherein said nucleic acid comprises at least one expression cassette comprising a gene which confers resistance to a selection agent.
- 7 45. The stably transformed duckweed plant according to Claim 44, wherein said gene which confers resistance to a selection agent is selected from the group consisting of *neo*, *bar*, *pat*, *ALS*, *HPH*, *HYG*, *EPSP* and *Hml*.
- 8 46. The stably transformed duckweed plant according to Claim 38, wherein said nucleic acid comprises two genes of interest.
- 30 9 47. The stably transformed duckweed plant according to Claim 38, wherein said nucleic acid encodes a protein or peptide selected from the group consisting of

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- 10 48. insulin, growth hormone, α -interferon, β -glucocerebrosidase, retinoblastoma protein, p53 protein, angiostatin, leptin, and serum albumin.
- 5 11 48. The stably transformed duckweed plant according to Claim 38, wherein said nucleic acid encodes at least one protein or peptide subunit of a multimeric protein selected from the group consisting of hemoglobin, collagen, P450 oxidase, and a monoclonal antibody.
- 10 12 62. The stably transformed duckweed plant according to Claim 38, wherein said nucleic acid encodes a secreted protein or peptide.
- 15 13 63. The stably transformed duckweed plant according to Claim 48, wherein said duckweed plant is from a species of *Lemna minor*.
- 14 64. A stably transformed duckweed plant tissue comprising a heterologous nucleic acid of interest incorporated in its genome.
- 20 15 65. The stably transformed duckweed plant tissue according to Claim 63, wherein said plant tissue is meristematic tissue.
- 13 66. The stably transformed duckweed plant tissue according to Claim 65, wherein said plant tissue is frond tissue.
- 25 16 66. The stably transformed duckweed plant tissue according to Claim 66, wherein said plant tissue is callus tissue.
- 17 67. The stably transformed duckweed plant tissue according to Claim 66, wherein said plant tissue is Type I callus tissue.
- 30 18 68. A duckweed tissue culture comprising the stably transformed duckweed plant tissue of Claim 63.

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69. A stably transformed duckweed cell comprising a heterologous nucleic acid of interest incorporated in its genome.
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70. 5 A stably transformed duckweed plant comprising a chimeric nucleic acid of interest incorporated in its genome, wherein said chimeric nucleic acid comprises a coding sequence operably linked to a transcription initiation region that is heterologous to said coding sequence.
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71. 10 The stably transformed duckweed plant according to Claim 70, wherein said chimeric nucleic acid comprises a duckweed coding sequence operably linked to a transcription initiation region that is heterologous to said coding sequence.
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72. 15 The stably transformed duckweed plant accordingly to Claim 70, wherein said chimeric nucleic acid is flanked by T-DNA border sequences.
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73. 20 The stably transformed duckweed plant according to Claim 70, wherein said duckweed plant comprises fewer than 5 copies of said chimeric nucleic acid.
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74. 25 The stably transformed duckweed plant according to Claim 70, wherein said duckweed plant is selected from the group consisting of the genus *Spirodela*, genus *Wolffia*, genus *Wolfiella*, and genus *Lemna*.
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75. 30 The stably transformed duckweed plant according to Claim 70, wherein said duckweed plant is selected from the genus *Lemna*.
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76. 35 The stably transformed duckweed plant according to Claim 70, wherein said duckweed plant is selected from the group consisting of a species of *Lemna minor*, a species of *Lemna miniscula*, and a species of *Lemna gibba*.
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77. 40 The stably transformed duckweed plant according to Claim 78, wherein said chimeric nucleic acid of interest comprises at least one expression cassette comprising a gene which confers resistance to a selection agent.

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28. The stably transformed duckweed plant according to Claim 27, wherein said gene which confers resistance to a selection agent is selected from the group consisting of *neo*, *bar*, *pat*, *ALS*, *HPH*, *HYG*, *EPSP* and *Hml*.
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- 5 29. The stably transformed duckweed plant according to Claim 28, wherein said chimeric nucleic acid comprises two genes of interest.
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- 10 30. The stably transformed duckweed plant according to Claim 28, wherein said chimeric nucleic acid encodes a protein or peptide selected from the group consisting of insulin, growth hormone, α -interferon, β -glucocerebrosidase, retinoblastoma protein, p53 protein, angiostatin, leptin, and serum albumin.
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- 15 31. The stably transformed duckweed plant according to Claim 28, wherein said chimeric nucleic acid encodes at least one protein or peptide subunit of a multimeric protein selected from the group consisting of hemoglobin, collagen, P450 oxidase, and a monoclonal antibody.
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- 20 32. The stably transformed duckweed plant according to Claim 28, wherein said chimeric nucleic acid encodes a secreted protein or peptide.
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- 25 33. The stably transformed duckweed plant according to Claim 28, wherein said duckweed plant is from a species of *Lemna minor*.
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- 25 34. A stably transformed duckweed plant tissue comprising a chimeric nucleic acid of interest incorporated in its genome, wherein said chimeric nucleic acid comprises a coding sequence operably linked to a transcription initiation region that is heterologous to said coding sequence.
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- 30 35. The stably transformed duckweed plant tissue according to Claim 34, wherein said plant tissue is meristematic tissue.
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- 30 36. The stably transformed duckweed plant tissue according to Claim 34, wherein said plant tissue is frond tissue.

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87. The stably transformed duckweed plant tissue according to Claim 84, wherein said plant tissue is callus tissue.
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88. The stably transformed duckweed plant tissue according to Claim 87, wherein said plant tissue is Type I callus tissue.
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89. A duckweed tissue culture comprising the stably transformed duckweed plant tissue of Claim 84.
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90. A stably transformed duckweed cell comprising a chimeric nucleic acid of interest incorporated in its genome, wherein said chimeric nucleic acid comprises a coding sequence operably linked to a transcription initiation region that is heterologous to said coding sequence.

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